

Supplemental Information

Functional Analysis of Orai1 Concatemers Supports a Hexameric Stoichiometry for the CRAC Channel

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SUPPORTING MATERIAL

Functional analysis of Orai1 concatemers supports a hexameric stoichiometry for the CRAC channel

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Table S1: PCR primers used to generate Orai1 subunit inserts

Primer	Purpose	Sequence
1	SU1 for	TCTAAGGCTCGAGGCCACCATGGAGCAAAAGCTCATTCTGAG GA
	SU1 rev	ACGGCAAGCTTGGCATAGTGGCTGCCGGCGT
2	SU2 for	TAATAAGCTTAAAACCTATACTTCCAAGGAGAGCAAAAGC TCATTCTGAGGA
	SU2 rev	ACGGCGAATTGGCATAGTGGCTGCCGGCGT
3	SU3 for	AGGAAGAACCGAGAACCTCTACTTCCAAGGAGAGCAAAAGC TCATTCTGAGGA
	SU3 rev	AGCAAGGTACCGGCATAGTGGCTGCCGGCGT
4	SU4 for	AGACACGGTACCGAAAACCTATACTTCCAAGGAGAGCAAAAG
	SU4 rev	ACGGCGACTGCAGCATAGTGGCTGCCGGCGTCAG
5	SU5 for	AATAAACTGCAGAAAACCTATACTTCCAAGGAGAGCAAAAGC TCATT
	SU5 rev	TACGGCGATATGGCATAGTGGCTGCCGGCGTCAGGG
6	SU6 for	TTACTAGATATCGAAAACCTATACTTCCAAGGAGAGCAAAAG
	SU6 rev	ATGCGTGGATCCCAGTGAGCTGCCGGCGT
7 (SDM)	ΔHind for	AGCCCGCCAAGCTCAAAGCCTCCAGCCG
	ΔHind rev	CGGCTGGAGGCTTGAGCTGGCGCG
8 (SDM)	TRS for	GGCCCGGGATCCACGAGAACCTTATTCCAGGGATCGGTCGC
	TRS rev	CACCATGGTGAG ACCATGGTGGCGACCGATCCCTGGAAATAAGGTTCTCGTGG ATCCCGGGCCCGC
9 (SDM)	L273D for	ACCGACAGTCCAGGAGGACAACGAGCTGGCGGAG
	L273D rev	CTCCGCCAGCTCGTTGTCCTCCTGGAACTGTCGGT

Primer sets were used for PCR amplification except where indicated. Primers used for site-directed mutagenesis are denoted by “SDM”

Table S2: Linking sequences in the Orai1 concatemeric hexamer

Orai1 Subunit #	N-term appended sequence	C-term appended sequence
1	XhoI-myc	HindIII
2	HindIII-linker ^a	EcoRI
3	EcoRI-linker ^a	KpnI
4	KpnI-linker ^a	PstI
5	PstI-linker ^a	EcoRV
6	EcoRV-linker ^a	BamHI
C-term eGFP	BamHI-TRS	

^alinker = TRS (ENLYFQG) + myc (EQKLISEEDL) + Gly-Ser stretch (NGGGGGGS)

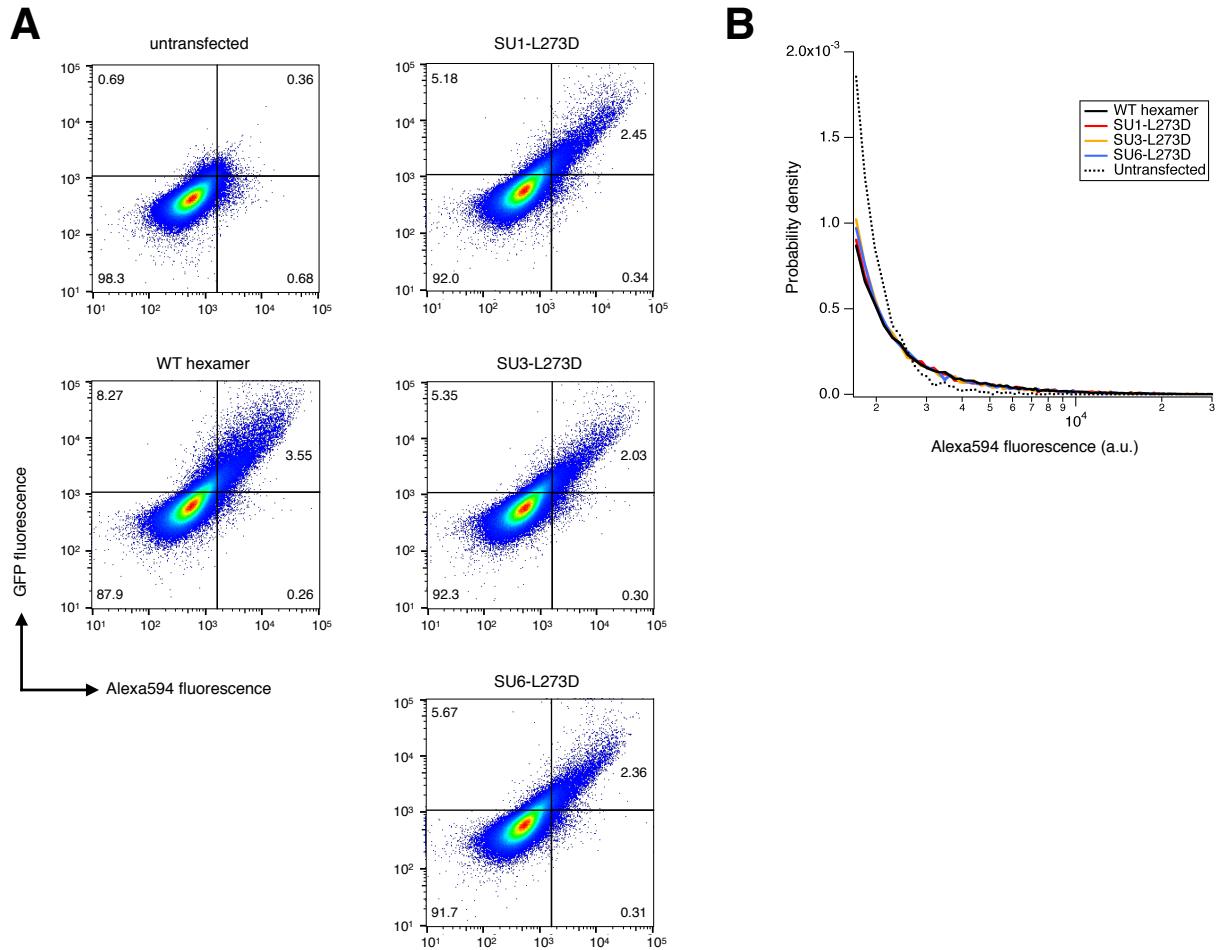


Figure S1. Flow cytometry analysis of 6xOrai1 expression in the plasma membrane. (A) Bivariate plot showing GFP (total Orai1) vs 2C1.1-Alexa594 (surface Orai1) fluorescence in untransfected HEK cells and HEK cells transfected with GFP-tagged 6xOrai1 variants. In cells transfected with 6xOrai1-GFP, GFP expression was proportional to Alexa594 fluorescence (upper right quadrants). Untransfected cells stained with 2C1.1 were used to establish a cutoff for GFP and Alexa594 negative and positive populations, as indicated by the quadrant gates. The gates were chosen separately such that 99% of untransfected cells fell in the negative population for GFP and for Alexa594. (B) Histogram of the GFP and Alexa594 double positive population in cells expressing Orai1 hexamer and untransfected cells. The Alexa594 fluorescence distribution is similar between WT and L273D hexamer populations. Each histogram contained > 1000 cells.